SEQUENCE LISTING

<110> Salmedix, Inc.
Leoni, Lorenzo M.

<120> COMPOSITIONS AND METHODS FOR THE DETECTION AND TREATMENT OF METHYLTHIOADENOSINE PHOSPHORYLASE DEFICIENT CANCERS

<130> 076936-0307942

<140> 10/779,476
<141> 2004-02-13

<150> 60/447,888
<151> 2003-02-14

<150> 60/460,715
<151> 2004-04-04

<160> 12

<170> PatentIn version 3.2

<210> 1
<211> 283
<212> PRT
<213> Homo sapiens

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Gly Thr Gly Leu Asp Asp Pro Glu Ile Leu Glu Gly Arg Thr Glu Lys
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Tyr Val Asp Thr Pro Phe Gly Lys Pro Ser Asp Ala Leu Ile Leu Gly
35 40 45

Lys Ile Lys Asn Val Asp Cys Ile Leu Leu Ala Arg His Gly Arg Gln
50 55 60

His Thr Ile Met Pro Ser Lys Val Asn Tyr Gln Ala Asn Ile Trp Ala
65 70 75 80

Leu Lys Glu Glu Gly Cys Thr His Val Ile Val Thr Thr Ala Cys Gly
85 90 95

Ser Leu Arg Glu Glu Ile Gln Pro Gly Asp Ile Val Ile Ile Asp Gln
100 105 110

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Phe Ile Asp Arg Thr Thr Met Arg Pro Gln Ser Phe Tyr Asp Gly Ser
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His Ser Cys Ala Arg Gly Val Cys His Ile Pro Met Ala Glu Pro Phe
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Cys Pro Lys Thr Arg Glu Val Leu Ile Glu Thr Ala Lys Lys Leu Gly
145                 150                 155                 160

Leu Arg Cys His Ser Lys Gly Thr Met Val Thr Ile Glu Gly Pro Arg
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Phe Ser Ser Arg Ala Glu Ser Phe Met Phe Arg Thr Trp Gly Ala Asp
            180                 185                 190

Val Ile Asn Met Thr Thr Val Pro Glu Val Val Leu Ala Lys Glu Ala
        195                 200                 205

Gly Ile Cys Tyr Ala Ser Ile Ala Met Ala Thr Asp Tyr Asp Cys Trp
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Lys Glu His Glu Glu Ala Val Ser Val Asp Arg Val Leu Lys Thr Leu
225                 230                 235                 240

Lys Glu Asn Ala Asn Lys Ala Lys Ser Leu Leu Leu Thr Thr Ile Pro
                245                 250                 255

Gln Ile Gly Ser Thr Glu Trp Ser Glu Thr Leu His Asn Leu Lys Asn
            260                 265                 270

Met Ala Gln Phe Ser Val Leu Leu Pro Arg His
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<212> DNA
<213> Homo sapiens

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catggcctct ggcaccacca ccaccgccgt gaagattgga ataattggtg gaacaggcct 180

ggatgatcca gaaattttag aaggaagaac tgaaaaatat gtggatactc catttggcaa 240

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aagtttaact atttaaaaga ctaaatgcac attttatggt atctgatatt ttaaaaagta 2160

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<211> 156
<212> PRT
<213> Homo sapiens

<400> 3

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Ala Thr Ala Ala Ala Arg Gly Arg Val Glu Glu Val Arg Ala Leu Leu
20 25 30

Glu Ala Gly Ala Leu Pro Asn Ala Pro Asn Ser Tyr Gly Arg Arg Pro
35 40 45

Ile Gln Val Met Met Met Gly Ser Ala Arg Val Ala Glu Leu Leu Leu
50 55 60

Leu His Gly Ala Glu Pro Asn Cys Ala Asp Pro Ala Thr Leu Thr Arg
65 70 75 80

Pro Val His Asp Ala Ala Arg Glu Gly Phe Leu Asp Thr Leu Val Val
85 90 95

Leu His Arg Ala Gly Ala Arg Leu Asp Val Arg Asp Ala Trp Gly Arg
100 105 110

Leu Pro Val Asp Leu Ala Glu Glu Leu Gly His Arg Asp Val Ala Arg
115 120 125

Tyr Leu Arg Ala Ala Ala Gly Gly Thr Arg Gly Ser Asn His Ala Arg
130 135 140

Ile Asp Ala Ala Glu Gly Pro Ser Asp Ile Pro Asp
145 150 155

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<211> 1218
<212> DNA
<213> Homo sapiens

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ggatttgagg gacagggtcg gaggggctc ttccgccagc accggaggaa gaaagaggag 180

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gccttcggct gactggctgg ccacggccgc ggcccggggt cgggtagagg aggtgcgggc 360

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gaagctgtcg acttcatgac aagcattttg tgaactaggg aagctcaggg gggttactgg 1140

cttctcttga gtcacactgc tagcaaatgg cagaaccaaa gctcaaataa aaataaaata 1200

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<210> 5
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<212> PRT
<213> Homo sapiens

<400> 5

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Glu Trp Arg Cys Ser Pro Leu Val Pro Lys Gly Gly Ala Ala Ala Ala
20 25 30

Glu Leu Gly Pro Gly Gly Gly Glu Asn Met Val Arg Arg Phe Leu Val
35 40 45

Thr Leu Arg Ile Arg Arg Ala Cys Gly Pro Pro Arg Val Arg Val Phe
50 55 60

Val Val His Ile Pro Arg Leu Thr Gly Glu Trp Ala Ala Pro Gly Ala
65 70 75 80

Pro Ala Ala Val Ala Leu Val Leu Met Leu Leu Arg Ser Gln Arg Leu
85 90 95

Gly Gln Gln Pro Leu Pro Arg Arg Pro Gly His Asp Asp Gly Gln Arg
100 105 110

Pro Ser Gly Gly Ala Ala Ala Ala Pro Arg Arg Gly Ala Gln Leu Arg
115 120 125

Arg Pro Arg His Ser His Pro Thr Arg Ala Arg Arg Cys Pro Gly Gly
130 135 140

Leu Pro Gly His Ala Gly Gly Ala Ala Pro Gly Arg Gly Ala Ala Gly
145 150 155 160

Arg Ala Arg Cys Leu Gly Pro Ser Ala Arg Gly Pro Gly
165 170

<210> 6
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<212> DNA
<213> Homo sapien

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gggcgcgctc agggaaggcg ggtgcgcgcc tgcggggcgg agatgggcag ggggcggtgc 180

gtgggtccca gtctgcagtt aaggggggcag gagtggcgct gctcacctct ggtgccaaag 240

ggcggcgcag cggctgccga gctcggccct ggaggcggcg agaacatggt gcgcaggttc 300

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cacatcccgc ggctcacggg ggagtgggca gcgccagggg cgcccgccgc tgtggccctc 420

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actcacgccc taagcgcaca ttcatgtggg catttcttgc gagcctcgca gcctccggaa 1140

gctgtcgact tcatgacaag cattttgtga actagggaag ctcaggggggg ttactggctt 1200

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<210> 7
<211> 105
<212> PRT
<213> Homo sapiens

<400> 7

Met Met Met Gly Ser Ala Arg Val Ala Glu Leu Leu Leu Leu His Gly
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20 25 30

Asp Ala Ala Arg Glu Gly Phe Leu Asp Thr Leu Val Val Leu His Arg
35 40 45

Ala Gly Ala Arg Leu Asp Val Arg Asp Ala Trp Gly Arg Leu Pro Val
50 55 60

Asp Leu Ala Glu Glu Leu Gly His Arg Asp Val Ala Arg Tyr Leu Arg
65 70 75 80

Ala Ala Ala Gly Gly Thr Arg Gly Ser Asn His Ala Arg Ile Asp Ala
85 90 95

Ala Glu Gly Pro Ser Asp Ile Pro Asp
100 105

<210> 8
<211> 897
<212> DNA
<213> Homo sapiens

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tctcacccga cccgtgcacg acgctgcccg ggagggcttc ctggacacgc tggtggtgct 240

gcaccgggcc ggggcgcggc tggacgtgcg cgatgcctgg ggccgtctgc ccgtggacct 300

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cagaggcagt aaccatgccc gcatagatgc cgcggaaggt ccctcagaca tccccgattg 420

aaagaaccag agaggctctg agaaacctcg ggaaacttag atcatcagtc accgaaggtc 480

ctacagggcc acaactgccc ccgccacaac ccaccccgct ttcgtagttt tcatttagaa 540

aatagagctt ttaaaaatgt cctgcctttt aacgtagata taagccttcc cccactaccg 600

taaatgtcca tttatatcat tttttatata ttcttataaa aatgtaaaaa agaaaaacac 660

cgcttctgcc ttttcactgt gttggagttt tctggagtga gcactcacgc cctaagcgca 720

cattcatgtg ggcatttctt gcgagcctcg cagcctccgg aagctgtcga cttcatgaca 780

agcattttgt gaactaggga agctcagggg ggttactggc ttctcttgag tcacactgct 840

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<210> 9
<211> 116
<212> PRT
<213> Homo sapiens

<400> 9

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Ala Thr Ala Ala Ala Arg Gly Arg Val Glu Glu Val Arg Ala Leu Leu
20 25 30

Glu Ala Gly Ala Leu Pro Asn Ala Pro Asn Ser Tyr Gly Arg Arg Pro
35 40 45

Ile Gln Val Gly Arg Arg Ser Ala Ala Gly Ala Gly Asp Gly Gly Arg
50 55 60

Leu Trp Arg Thr Lys Phe Ala Gly Glu Leu Glu Ser Gly Ser Ala Ser
65 70 75 80

Ile Leu Arg Lys Lys Gly Arg Leu Pro Gly Glu Phe Ser Glu Gly Val
85 90 95

Cys Asn His Arg Pro Pro Pro Gly Asp Ala Leu Gly Ala Trp Glu Thr
100 105 110

Lys Glu Glu Glu
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<210> 10
<211> 1515
<212> DNA
<213> Homo sapiens

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ttcattcatt cactc 1515

<210> 11
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer for cloning MTAP cDNA

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<210> 12
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Antisense primer for cloning MTAP cDNA

<400> 12
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